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TITLE: ANALYSIS OF DENGUE VIRUS ENHANCING EPITOPES USING

PEPTIDE ANTIGENS DERIVED FROM THE ENVELOPE GLYCOPROTEIN

GENE SEQUENCE

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The biological events leading to the development of severe disease manifestations of dengue infections (dengue hemorrhagic fever/dengue shock syndrome, DHF/DSS) has been the focus of many investigations. Though DHF/DSS cases occur during primary dengue infections, the majority of severe dengue disease has been associated with persons experiencing a secondary heterotypic dengue infection. What do we know about the pathogenic processes that result in the exacerbation of dengue symptoms? Can these factors be identified in vitro and then correlated with clinical disease? The results of previous studies suggest that pre-existing DEN antibodies circulating in a person may be a risk factor in the development of DHF/DSS. However, other host and viral factors must be involved since only a small proprotion of the population residing in endemic dengue disease regions develop severe dengue disease.

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FOREWORD

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TABLE OF CONTENTS

REPORT DOCUMENTATION PAGE	i
FOREWORD	i
TABLE OF CONTENTS	i
	1
	2
	2
Reactivities of anti-DEN-2 E glycoprotein synthetic peptide antibodies with DEN-1 and DEN-2 virus Cloning and sequencing the nucleotides encoding the	2
non-structural DEN-1 CV1636/77 genes	3 3
Relevance of epitope mapping using anti-peptide	4
Analyses of the genetic relatedness of DEN viruses	4 4 5
REFERENCES	6
APPENDIX 1	s
APPENDIX 2	s
APPENDIX 3	'S
APPENDIX 4	

INTRODUCTION

The biological events leading to the development of severe disease manifestations of dengue infections (dengue hemorrhagic fever/dengue shock syndrome, DHF/DSS) has been the focus of many investigations (1,8). Though DHF/DSS cases occur during primary dengue infections, the majority of severe dengue disease has been associated with persons experiencing a secondary heterotypic dengue infection (6,7). What do we know about the pathogenic processes that result in the exacerbation of dengue symptoms? Can these factors be identified in vitro and then correlated with clinical disease? The results of previous studies suggest that pre-existing DEN antibodies circulating in a person may be a risk factor in the development of DHF/DSS (9). However, other host and viral factors must be involved since only a small proportion of the population residing in endemic denque disease regions develop severe denque disease.

Antibody-dependent enhancement (ADE) of dengue virus (DEN) replication in vitro has been proposed as a pathogenic mechanism in the development of DHF/DSS. We believe that enhanced virus replication is an important part of DEN pathogenesis and that ADE is the result of the interplay of host cells, antibody and virus strain. We investigated DEN virus replication in order to define the parameters that could lead to enhanced growth in the presence of antibodies to flaviviruses.

SUMMARY OF PREVIOUS ACCOMPLISHMENTS FOR FY 1990

- Developed a sensitive ADE assay to examine the interplay of antibody, virus and host cells in DEN replication. Human promonocytic HL-CZ cells supported enhanced DEN replication better than K562 or U937 human cells at low MOIs (10⁴-10⁶).
- Determined that the sensitivity of the HL-CZ cells was related to the number of Fc-receptors (FcRs) and the FcR types (FcR II > FcR III > FcR I) expressed on the cell surface since blocking the FcRs by specific FcR monoclonal antibodies (MAbs) abrogated virus replication.
- Characterized the abilities of MAbs 4G2 and 3H5 to mediate virus neutralization and enhanced replication. Determined that the MAbs recognizing DEN envelope (E) glycoprotein epitopes were involved in NT and ADE and confirmed the observation that these activities were concentration related rather than epitope-directed.
- Compared the growth profiles of selected DEN viruses representing epidemiologically important genotypes using our ADE system. Viruses from endemic dengue region (Thailand) were more antibody-dependent than that of viruses from the dengue epidemic region (Caribbean).

- Attempted to define the antigenic epitopes that mediate ADE and NT by using antibodies raised against DEN-2 E glycoprotein synthetic peptides. We were able to determine that some antipeptide antibody pools were reactive in ADE and/or NT tests, though reactions appeared much weaker than the MAb controls.
- Continued to determine the nucleotide sequence encoding the non-structural (NS) regions of DEN-1 CV1636/77 virus. Of the regions examined thus far, it appeared that DEN-4 Dominica and DEN-1 CV1636/77 were very similar in the NS region, suggesting that the viruses have evolved closely or that there may be genomic recombination.

ACCOMPLISHEMENTS FOR FY 1991

Examination of FcR-expression in HL-CZ promonocytic and enhanced virus replication. Dr. Wu Tse Liu of the National Yang Ming Medical School, Taipei, Taiwan provided us with several lines of HL-CZ cells (14). We compared the virus yields of the uncloned, clone 3 and clone CCC-5 cells. The results of the enhanced growth profile of DEN-2 16681 virus were variable thus suggesting that the permissiveness of the cells to support enhanced virus replication could be intrinsically different. We investigated the discrepancy of virus yield of these cultures and determined that if the ADE results were grouped by >20% FcR-expression (rosetting) compared with virus yield (>10' pfu/ml), the presence of rosetting cells was a factor in cell permissiveness (Table 1) as we had demonstrated before when comparing virus growth in U937, K562 or HuPBL cells. However, in a few cases, neither high nor low number of rosettes correlated with virus yield, thus implying that some other factors also affect virus replication. Therefore, in our analyses of ADE results, we only accepted the tests in which 1) virus control cultures (no antibody) contained no visible growth and 2) positive ADE controls of cultures containing antibody produced 10' pfu/ml virus yield.

Reactivities of anti-DEN-2 E glycoprotein synthetic peptide antibodies with DEN-1 and DEN-2 virus. We have previously determined the reactivities of mouse antibodies immunized with DEN-2 E glycoprotein peptides in PRNT and ADE assays. Though we were able to demonstrate some activity in sera obtained from mice immunized by i.m. and s.c. routes, there was variation between antibody specimens directed against the same peptide heterologous mouse strains (19). To normalize antibody reactivity, sarcoma-primed BALB/C mice were immunized with individual DEN-2 E glycoprotein synthetic peptides and the resulting mouse ascites fluids (MAFs) were assessed (Table 2). To compare the anti-peptide antibodies on an equal basis, immune mouse ascites fluids (MAFs) were purified by elution from protein A columns and standardized to 1 mg/ml and examined by ELISA. The reactivities of the purified Igs were not appreciably different from crude MAFs suggesting that binding to virus antigen or peptide was related to the original strength of the MAF and was not affected by increased/decreased affinity of the antibody to the peptide.

In addition to testing each of the anti-peptide antibodies individually, we attempted to reconstitute antigenic binding sites by mixing the antibodies together (Table 3). Seven anti-peptide antibodies were mixed in various combinations and each tested for binding to antigen, for ability to neutralize virus and for mediating enhanced virus growth. Binding of MAFs, 1 μ g and 10 μ g of anti-peptide antibodies to DEN-2 Jamaica virus antigen were examined by ELISA, and these results reflect original MAF titers. HL-CZ cells were infected with DEN-1 16681 virus at MOI of 10°. Where the positive ADE and background culture controls cells met our acceptance criteria, the enhanced virus yields in the presence of 1 µq of anti-peptide antibodies was not consistent (Table 3). Neutralization activity of the anti-peptide antibodies DEN-2 16681 virus was examined. One μq of an antibody was mixed with 1000 pfu of virus and incubated at 37 C for 1 hour. surviving virus was ascertained by testing the virus/Ab mixes in the BHK-21 plaque assay (16) and the results reported as percentage of reduction of input virus (Table 3). Both anti-peptide mix 21 (antibodies to peptides 06, 240, 274, 17 and 361) and mix 22 (antipeptides 240, 274, 17, and 361) neutralized 53% of the input virus, and anti-peptide mix 42 (antibodies to peptides 240 and 274) neutralized 100% of the input virus.

Cloning and sequencing the nucleotides encoding the non-structural DEN-1 CV1636/77 genes. The genome encoding the structural genes of DEN-1 CV1636/77 have been previously published (3). Last year we completed the nucleotide sequences encompassing NS4a, NS4b, and NS5 genomic regions of DEN-1 CV1636/77 and compared the results with the published sequences of DEN-2 (5), DEN-3 (18), and DEN-4 (15). We have completed the sequencing of the rest of the non-structural nucleotide regions for NS1, NS2a, NS2b, NS3 and the final 800 base pair sequence of the 3'-end of the viral RNA (Figure 1).

The similarity of the non-structural gene regions between the 4 DEN viruses are summarized in Figure 2. DEN-1, DEN-2, and DEN-3 share between 63%-68% similarity by nucleotide sequence comparison and 65%-78% similarity by their deduced amino acid sequences.over each of the non-structural gene regions. DEN-1 and DEN-4 however are very similar in NS3, NS4a, and NS4b regions (95%-98%); this is contrasted by a comparison of DEN-1 and DEN-4 over the NS5 region where similarity extends to only 78%. When the combined nucleotide and deduced amino acid sequences over NS3, NS4a, NS4b, and NS5 regions are examined, the similarity of 90% remains between DEN-1 and DEN-4 (Figure 3,4).

DISCUSSION

Studies to identify genomic correlates of DHF/DSS have primarily depended mouse, monkey and other laboratory experiments. Animals infected with DEN will respond by seroconversion to the infected virus but do not develop DHF/DSS. Therefore most disease parameters of infected humans are based on clinical observations, and epidemiological assements. The availability of genomic sequences of flavi- and dengue viruses, the synthesis of peptides corresponding to deduced amino acid sequences, and the development of molecular techniques, make it possible to begin studies to correlate biological activities with genomic variation.

Relevance of epitope mapping using anti-peptide antibodies. the standardized ADE assay, we have been able to examine divergent aspects of the roles of antibody, virus and host cells. From these studies, we could determine that each component share roles in virus replication and that the ADE system developed could isolate many of these factors for further examination. In our attempts to identify the genomic regions that encode antigenic epitopes involved in biological responses using anti-peptide antibodies, we could identify the epitopes that mediate neutralization of the virus. However, the enhancing activity could not be located within a specific region using the anti-peptide antibodies. This does not mean ADE activity cannot be found, the non-or broad reactivity may only be a matter of using sub-optimal antibody concentrations, the the anti-peptides, inappropriate mix of the virus variability or the susceptibility of the host cell stage to virus replication. It is likely that in using anti-peptides to map strategic epitopes, a reactive result implies reconstitution of epitopes, whereas non-reactivity implies only that epitopes were not reconstituted.

Using anti-peptide sera directed against selected DEN-2 E glycoprotein regions, we have been able to demonstrate that some of the E regions will elicit antibody that will mediate enhanced virus replication, neutralization and enhancement, and neutralization alone. Though this series of experiments need to be repeated with other antibody mixes and other DEN viruses, the neutralization epitope defined by amino acids 240-274 is is the first direct association of specific E-glycoprotein regions with biological functions.

Analyses of the genetic relatedness of DEN viruses. The similarity of the non-structural genomic sequences of DEN-1 and DEN-4 viruses was unexpected. Published sequences of NS regions of DEN-2 and DEN-3 viruses led us to expect only 63%-68% homology among the DEN viruses. DEN-1 and DEN-4 viruses, however, appeared to be nearly identical from NS1 through NS5 and again at the 3'-end, suggesting that there may be a double-crossover genetic recombination. To preclude that possibility that the DEN stock cultures may have been contaminated with one or the other viruses, we plaque-picked purified virus from our CV1636/77 stock and found that they all

contained the same characteristics (data not shown). In addition, we are examining, by PCR amplification and sequencing, of other DEN-1 and DEN-4 strains in the 3'-end region to analyze the relationship.

Future studies. During this study, we were not able to achieve all the goals set forth in the original proposal. We would liked to have been able to get anti-DEN-1-peptide serum to analyze and work as we did with DEN-2 peptides. However, we were delayed by the unexpected complication of the DEN-1 sequence homology with DEN-4 and felt that was important finding to follow-up and analyze. We are planning to do work on examining patient serums that would recognize DEN-2 peptide mix 240/274 and to identify whether this region of the E-glycoprotein is a useful diagnostic tool.

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Tables

- Table 1 Chi-square comparison of the relationship between rosette expression and virus replication in HL-CZ cells
- Table 2 Reactivity of the mouse ascites fluids obtained from mice immunized with DEN-2 E glycoprotein synthetic peptides
- Table 3 Mixtures of purified anti-DEN-2 synthetic peptide immunoglobulins and PRNT/ADE results

Figures

- Figure 1 Location of CV1636/77 clones sequenced and the diagram of the similar regions between DEN-1 and DEN-4.
- Figure 2 Percentage of homology of the non-structural region nucleotide and deduced amino acid sequences of DEN viruses.
- Figure 3 Comparison of the nucleotide sequences comprising the non-structural genome regions of DEN-1, DEN-2, DEN-3 and DEN-4 viruses.
- Figure 4 Comparison of the deduced amino acid sequences of the non-structural regions of the four DEN viruses.

TABLE 1
Chi-square comparison of the relationship between rosette expression and virus replication in HL-CZ cells

PFU VIRUS/% ROSETTE	> 20% ROSETTE	< 20% ROSETTE
> 103 VIRUS/ML	6*	1
< 10 ³ VIRUS/ML	1	4

^{*}p <0.001

TABLE 2 REACTIVITY OF ANTI-PEPTIDE ANTIBODIES AGAINST HOMOLOGOUS PEPTIDE, DENGUE-2 AND DENGUE-1 VIRUSES

PEPTIDE"	ANT: PEPTIDE	IGEN USED IN E	LISA ^b DENGUE-1
1-2*	6400	1280	320
35	102400	128000	4000
3-8/1	100	<10	<10
4-6	400	40	10
04*	1600	160	10
142*	3200	640	10
167*	100	10	<10
06*	25600	5120	40
240#	1600	1280	80
274*	100	10	<10
16*	800	160	80
17#	3200	160	<10
361*	12800	640	80
437*	3200	20	80
+control		>12800	>12800
-control		<10	<10

[&]quot; Roehrig et al (19).

Each well of the ELISA plates were coated with either 1 ug of peptide, or gradient-purified virus antigen. The antibody titers as reported as positive were adjusted by subtracting 2 standard-deviations above background reactions.

^{*} Synthetic peptides which are distinct by deduced amino acid sequence from DEN-2 Jamaica.

[#] Synthetic peptides that are not distinct from DEN-2 Jamaica virus (>90% homology) but are in the NT/ADE region mapped by anti-peptide antibodies.

TABLE 3 ANTIBODY-DEPENDENT ENHANCEMENT (ADE) AND PLAQUE REDUCTION NEUTRALIZATION (PRNT) RESULTS

	ANT				BODIE		-		PR	ADE ^c	
1-2	35 3	3/8-1	4 - 6	06	240	274	17	361	DEN1	DEN2	pfu/ml
									20	35	250
									15	41	< 7
		: .							19	32	< 7
	_								24	35	< 7
									15	32	>8300
									24	31	280
						M:			31	32	4600
									27	36	>8300
	_								28	46	< 7
									32	38	< 7
									36	42	3700
									42	56	260
									36	42	1400
									22	53	400
									18	53	< 7
									38	40	< 7
									10	19	< 7
									28	54	< 7
									16	56	< 7
									4	38	1500
									8	35	4000
									4	32	4500
									35	100	800
									16	27	5500
									0	10	3600
4G2	MAb*	, fla	vivi	rus	group	reac	tive		96	9 7	>8300
3H5	MAb*	, den	igue - :	2 spe	ecifi	C			12	100	8300
Viru	s co	ntrol	.@						0	0	< 7

Legend for TABLE 3

Purified anti-DEN-2 peptide antibodies, used 100 ng of each anti-peptide in the mixes. Filled boxes indicate the antipeptide serum used.

PRNT by BHK-21 semi-micro assay. Average plaques of DEN virus in each 24-well = 26. Serum and virus were incubated at 37 C for 1 hour. Percentage of neutralization shown here. >50 % reduction shown in **bold face**.

ADE assay with HL-CZ cells and MOI of 10.5. Cells were infected and incubated for 4 days at 37 C in 5% CO₂. Virus replication was assessed by plaque titration in BHK-21 semi-micro assay. Results are the average of 3 experiments.

* The amount of 4G2 used for PRNT = 10 ug, for ADE = 1 ug. 3H5 used for PRNT = 1 ug, for ADE = 100 ng.

• Virus control = only virus without antibody for PRNT and ADE.

ABSTRACT

ASTMH 1991

THE NON-STRUCTURAL GENOME OF DENGUE-1 VIRUS CV1636/77: COMPARISON OF THE NUCLEOTIDE AND DEDUCED AMINO ACID SEQUENCES WITH THE OTHER DENGUE SEROTYPES.

Chu MC, Putvatana R, and Trent DW. Division of Vector-Borne Infectious Diseases, National Center for Infectious Diseases, Centers for Disease Control, Ft. Collins, Colorado 80526

The determination of the entire nucleotide and deduced amino acid sequences of the DEN-1 Jamaica CV1636/77 virus together with the known sequences of DEN-2, DEN-3 and DEN-4 viruses provide a complete sequence database of the DEN serotype We have previously presented analyses of the complex. nucleotide sequences encoding the structural and the nonstructural NS3-NS4-NS5 regions. Sequence analyses of the entire genomic RNA and cloned cDNA of DEN-1 reveal that the genomic RNA contains 10,641 nucleotides and encodes an open reading frame of 10,224 nucleotides that translate into 3408 amino acid residues. A comparison of the DEN-1 sequence with DEN-2 (Jamaica; Deubel et al., 1988), and DEN-3 (H87; Osatomi et al., 1990) reveal that the genomic sequences encoding the structural (C-prM-M-E) and some of the non-structural (NS1, NS3, and NS5) regions are conserved (>70% homology). 3'non-coding regions of DEN-1, DEN-2, DEN-3, and DEN-4 (Dominica; Mackow et al., 1987) are of different nucleotide lengths comprising 387, 455, 448, and 385 base pairs respectively. The entire non-structural sequences for DEN-1 and DEN-4 share >90% identity. These results reveal a close evolutionary relationship between Caribbean DEN-1 and DEN-4 viruses which cannot be detected by conventional serological methods. Thus the sequence database of all the DEN virus serocomplex will be a useful tool in studying the genetic variation and evolution of these viruses.

ABSTRACT

dengue (DEN) enhancement Antibody-dependent (ADE) of infection in human mononuclear cells in vitro has been standardized using a human promonocytic cell line HL-CZ, purified monoclonal antibodies (MAbs), and select DEN viruses. Characterization of the Fc-receptors (FcRs) expressed on HL-CZ cells have indicated that subsets of FcR mediate ADE better than others. Using this standardized system, we have compared the ability of mouse anti-DEN 2 envelope (E) peptides to elicit virus neutralization and ADE. Peptides 1-2, 437 appear to elicit ADE activity in contrast to other peptides that appear to elicit neutralization but not ADE. Though these assays need to be repeated, it appears that differential functions may be attributed to particular E genomic regions. The comparison of the nucleotide sequences of DEN-1 RNA encoding the non-structural proteins to the other DEN sequences has revealed that DEN-4 and DEN-1 share > 90% similarity in NS3 and NS4a, 4b genome regions. DEN-3 and DEN-1 have a deletion in in NS5 that is conserved in other DEN-1 and DEN-3 isolates. These genomic comparisons indicate that non-structural differences need to be studied as well for our understanding of DEN replication and pathogenesis.

PUBLICATIONS

- Chu MC, Huang GH, Collins ND, and Trent DW. An experimental model for analyzing antibody-dependent enhanced growth of dengue virus in human promonocytic HL-CZ cells. Submitted for publication.
- 2. Chu MC, Putvatana R, Huang GH, Roehrig JT, and Trent DW. Analyses of the reactivities of anti-dengue-2-E glycoprotein antibodies to dengue-1 and dengue-2 viruses. Manuscript in preparation.
- 3. Chu MC, Putvatana R, Repik P and Trent DW. Similarities of the dengue-1 CV1636/77 and dengue-4 Caribbean viruses: genomic sequence comparison and oligonucleotide fingerprinting. Manuscript in preparation.

DENGUE VIRUS GENOME ORGANIZATION NON-STRUCTURAL REGIONS NS5 OVERLAPPING CDNA CLONES 4a,4b NS4 NS3 2a,2b NS2 STRUCTURAL NS1 5

The second secon

REGIONS OF > 90% IDENTITY BETWEEN DEN-1 AND DEN-4

(MARKED CLOSED AREAS BETWEEN THE ARROWS)

§ 3.END NS5 NS4 NS3 NS2 NS1

Fisher 1

COMPARISON OF THE NON-STRUCTURAL GENOME OF DENGUE-1 CV1636/77 WITH DEN-2, DEN-3, AND DEN-4

	DEN-2	2-2	DEN-3	ლ	DE	DEN-4
	LZ	AA	Ę	AA	Ę	AA
NS1	%89	%69	%69	29%	%26	84%
NS2a	36%	34%	38%	39%	%66	%86
NS2B	%69	23%	29%	22%	%86	%26
NS3	%69	71%	%02	74%	%86	%26
NS4a	64%	%89	61%	%59	%66	%26
NS4B	%89	83%	%69	85%	%86	%26
NS5	%02	%92	71%	81%	73%	%62
3'-END	%29		72%		%26	

FIGURE 2

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COMPARISON OF THE NS1 NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

non n		A AAA		. GCG		_	A	•		S NGG			:				უ.				ع : :				A CA		
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COMPARISON OF THE NS2a NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

GGC ACA UCA GAA ACU UUU UCU AUG GGU CUG UUG UGC CUG ACC UUG UUU GUG GAAG CAG AUUC .ACA C.AA G.C G.A A G.A CC CA .AG GUGC .ACC A.A G.CU U G.A A.C C.C U.U	ACU AGG AAA CAC AUG AUA UUA GUU GUG GUG GGA .CU GCA C.G C.AU .CA GGG .AAU GCGGU C.C	CGA GCC CUC AUC AUG AUG A.G G.U AC A.A A	AUC AUG GCA GUG GC. C.A CUCA GCU C.A AUU .CA	GUG CUG GGU GUG UUU UUA AGG AAA CUC ACU UCA AGA GAG U.U GCA .CGA C.A C.C UUG. AAA CUG ACC UCC A U GCU UUG .GACC CUGG AAA CUG AC. UCU AGA	AUG GUA AUA GGA AUG GCC AUG ACACC .CC AU. GGA AUC GCA CUC U C.G GGUU GG. UUG GCC .UG	AUU GAU GGA AUA UCA CUG GGA CUA AUU UUG CUA AAA AUA GUA ACA CAG UUU GAC C.G ACAU GCG .U. GCC CUG GGCG A G.C CTC .A. A GUG AGA AA. AUGG .CG AAU GG. AUU GCU UUG GGG C.C A GCU CUU .A. C.G .U. ACA CAA UUU AG
CAG	UGC	3604 UGU GCU A GUG A.A U GUG CUC C	GAC	AUG	CUA	GAA
u	AUG		. CU	. AA	A.G	GAA
AGU	GUG		UC.		U	. AA
DEN-1	DEN-1	DEN-1	DEN-1	DEN - 1	DEN - 1	DEN-1
DEN-2	DEN-2	DEN-2	DEN-2	DEN - 2	DEN - 2	DEN-2
DEN-3	DEN-3	DEN-3	DEN-3	DEN - 3	DEN - 3	DEN-3
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COMPARISON OF THE NS2B NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

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		C:C			GNG	:	:	:		CUA	ტ		:		AUC	:	. AU	:		AGU	CAA	.AC	A.		GCA							
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COMPARISON OF THE NS3 NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

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	CNG		:		GUA	ე ე	D	:		AUC	ე ე	U.G	GAU		UCA	:	:	:		CNC	ტ	A.T	:		ACC	:	Α	:	ָ ֖֖֖֖֖֓ ֖֖֖֓֞	۸ (:	:
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	ACU	9 ⋖	•		AAA	U.C	:	:		ACA	:	:	:		AGG	.AG	. AA	:		GAA	ט	r	:		CCO	:	A	:	ر د د	5	•	:
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(a) AAA GCA CUC UAU GCA AAU GCA GUA GUU ACC AAA UCA GCA GCA GCA GCA CAA C	CGA AAG AAA AGA UUA ACU AUA AUG GAC GCU AA CGAU CC	CUU CCA UCA AUA GUG AGA GAA GCC UUA AAA G.CU A G.UUU A G.UUU A	ACGU	ACC CCA GCU GUG A C A.C A A.UA ACA	CONTRACTOR ACA ACA AGA CUU UUG UCA UCA ACC AGG GUU CCA AAU UAC AAC CONA U	JG GAU GAA GCA CAU UUC ACC GAU CCC UCU AGU GUC GCG GCU AGA GGA UACC CC ACA G.A AAGU ACA G.C A.A
AAA GGA A G.G AGU GCC A	UUU C AUGC A	AGA AUU C UACA. UA	GCU CCC A	UAU CAGCA	CAU GCA A	GUG AUG A A.A
AAC AGG GA A 5008 UAC GUC U	5068 GAC AUU GAG	5125 ACA AAA G CGG	5185 AUU UUAC C.G	5245 AUC CGU A A.A A A.G	AUG UGU	5365 CUU AUA GC U.G
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		ACU G	UCG GAAU .		AGA G		AGA G
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40C	GAA AUG C	GAC U	AAU GC. C	GAG	A UG	AUC A	AGCU
AUCU	AUA C U	ACA G u	GCA	ACA U.U 	GAA	GUU G	GCA U.U G
900 900 	CCA	AUA G.C u	AUU	GAU C	UCU A	CCA G	CCA .AC GU.
	AGC GCA GCU				AUA U		ACUC
GCA	AA CU	GACA	AA U	ACC U	GAC	CUC A.G	GUG
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UCG A	GCG AGU A.A	GAA		AAA G	GAC u	AGA G	GUC G
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DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4
	дддд				пппп		

G AAC CCA GCA CAA GAC GAC CAA UAC GUU UUC UCC GGA GAC CCA CUAA .U AA. A.U G A.A .A. AUGGAU G A.A .A. AUGGAU G A.A .A.A .	GAU CAU GCC CAC UGG ACA GAA GCA AAG UG A A A	AUC AUU CCA ACA UUG UUU GGU CCG GAA AGG GAA AAA ACC CAA GCC	CGC CUC AGA GGG GAA CAA AGG U U.G A GC GC GC	GUG UGG CUG AGC UAU AAG GUA GCU UCU GCU GGC AUU UCU UAC AAA C C GC GC GA G AA C AAC GC U A GC. C A A A A A GC C AA C GC GC A	UUC ACA GGG GAA AGA AAUu GAUa AUU .AGCu GAUA C.U	AGA GAG GGA GAA AAG AAA AAG CUA AGG .AAGGGA UAA .AGA	CCC AUG GCU UUG AAG GAU UUC AAG GAG UUU GCC AGU GGA AGGA CA C.AAAA GC GAAU U.AA C.CA
AUA GGA AGG	AAU GAU GAA GA!	ACC CCA GAA GGG AUCAUAA	GAG UUU A .AC A .AC	3AC UUC CCC A	CGG GAA UGG UGC UUC A.A AGGUU A.A A	AUU UGG ACUCACA	UAC GCU GAC CC
DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4

COMPARISON OF THE NS4a NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

AGG . A.	0	UUA C.C A C.G	AAAG	A AUU	AUA	A G
UCU CAG		GCU ACA . GA	GGG	GAA C.G	0.0 0.0 0.0	UUG A.A A.A
UCC A.U G	GGG 	GUA C.G C.G	AUA U	GCA	CUG U	A U A G.C G.G
CUU ATG U.A	GGA	CUU U.A 	GGA u	GUA UAU A.G	GUA g	GUC
UAC .U. C	AGA GC. CAU	AUG C.U U.A	AAA 	UGG	A UG A 	UAC u
ACU UCA	GAA	cuc g	GGG ••• A ••• U	CUC	CGC .U. AUG	AUC .C. GCA
CCA	ACA G.U U	ACA	CAA AGC UC.	UUG C.C A	DDD	000 0.0
UUG C.A G	ACAG	GAA G	AUG	GGC AU. 	DDDD	CAA
AGU G	CAC	CUG	UUCA	AGU	GAG	AA U
900 90.	CUC	UCA A.C A.G	B	GCU U.C	CUA G	GAC
AUUG	AUG G	GAG	CUG U.A	GUG AC. . CU	AUA G.C	CAA
GAG A	5 00.0 00.0	CCG	UUC A.G	GCG AUC AUU	A UC A	CCA
ACA	AUA U U.G	CUU G	AUC GCA	AUU A G.A	UCA G.U	ACCA
CUA AUC G.G	AAC	GAA	GGU A	ACC UG. UGU	GCC u TCG	AGG A A
AUC C.A C.U	GAU	AAC .GU G.G	GCA .g.	AUA UGU U	GCG A	CAA G
GAC A	CUU CUU CUU	CUG C G	ACA (UUG A C.C	AUA	AAA G
cuc 	GCC	GCC u	AUG G U.A	GGU A	UGG	GAA
ACUC	CUC GA. AA.	CAC u u	GCU A.A UUG	AUG C	CAG C	CC
AUA U.G	AAG GA .GA	CUU AA.	GGU . CC AUC	UCA A.C	CCC	GAA
NS4a 6382 AGU A UC. UC.	6442 GCC 7 A.G	UAU UAU O C	CUA CUA A.G. A.G.	A ACU	CAA	000 000 4 · · ·
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 3	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 3

6831 DEN-1 AUU CUC ACC AUC AUU GGU CUA AUA GCA GCC DEN-2 A G.G G.G .CC GCCC AUGA DEN-3 AUA U.G GCCA A GG DEN-4	6831	GCC	Α:	:	:
	•	GCA	AUG	r C	:
		AUA	ე.	: :	:
		CUA	ga.	Α	:
		GGN	ეე.	.CA	:
		AUU	G.G	gc.	:
		AUC	G.G	U.G	:
		ACC	A :	Α	:
		CUC	•	\mathbf{n} .	:
DEN-1 DEN-2 DEN-3 DEN-4	6802	AUU	ບ:	Α	•
		DEN-1	DEN-2	DEN-3	DEN-4

COMPARISON OF THE NS4B NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

A ACA U . GGU	G CUC	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	A AAA	u ucu c	A AUA CUU	U GGG AC A
7 . C	ACG	ACG U.C U.C	J GGA G	UAU 	J GCA C	J GCU
GUA A.U .A.	NGG	AAC u	CUU U.A	100 100 100 100 100 100 100 100 100 100	U A U 	GCU
CAG AGC A.A	GCA	GAA	666 0 U	GGAU	CAU	ACA G
UAC GGA . CU	UCA	AUA U	AUG	AUG C	GUC CA ACA	AGG A
UUU G A.G	GCU	ACC . G.	CUA	GCA 	G.C.	AAA
GGG	CCA	CAC U	GUCG	UUA C.U G	CUU U.G 	CAG
0.0 0.0 A	AGA C.U CAC	AGA	GCC A.A .UG	CUG	AUG U.A U	GCC u
GAU C	UUG C.A 	CUG U U	GCA 	000 0 · · · O 0 · · · · · · · · · · · · · · · · · · ·	GGC CUU CUU .U.	GAG A
ACG . AA GA	GACU	AUG	CAG A	GUG A	UUA C.U G	AGA C.U
AAA G	GUG A.A	CCC	AA C	GGU A	UCC G.U G.A	ACA C U
ACA	GAU C	ACUA	GCC u	CUC A U.G	GCAG	GCC A
AAA 	cuc g ug	CUG G.U A.A	A UU 	GAC C	ACA	AAA
GAA	AUC	AUU U G.A	3CC	A UGC	UUG C.C C.U	GCA
AUU C.G U.G	ACC	A C A	GCA A	AGA . AG . A.	ACC u u	CAG
cug u.c	ACC .G.	ACC	CUA	CAC UCA UCG	ACA .U. CU.	UUG C.U
GGG 	GAA G CC.	GCC 	 	CUC U.G A.A	CCA u	GGA
AUG	ucu	GUA G	CUA G.G G.G	CCG	AA C	CC.
GAG	GAA	80.0 0	AACu	UGG	300 300 300 300 300 300 300 300 300 300	GGC G
MS4b 6829 AAC 6889	CAG	UAU UAU O O C	GCC 7. UG A	GGA UGG	CAA GUGGGC	AUA
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

DEN-1 DEN-2	AUC AUG AAA AAU CCC ACA	AAA	AAU C	CCC	ACA	gug c	GAC (GGG 1	AUA A	ACA (GUA 1	AUA GAU	3AU o	CUA (GAA (CCA	AUA I	UCC T	UAU
DEN - 4	7300	ง . : :		₹ :) : : :			£ :			٠. د ٠. د		٠ . • •						
DEN-1	GAC CCA	AA	A UUU GAA AA	GAA	Ŋ			995		SUC 1			CUA	one of) DQC		GCU (AA
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DEN - 4	7360	•	•	:	•			:	:	•			:		•	•	:		:
DEN-1	CUA CUC	JUG	AUG	AGA	ACA	ACA		3cn 1		JGU (oug.	ACU 1	JUG (၁၁၅	ACA (CA CA
DEN - 2 DEN - 3	G.: U.A	۲.	•	<u>ა</u>	D :			ָ ניי		•		D	C. A	ບ ບ :	A A	U	υ ·	o :	n :
DEN-4	74.0				· ·	:		•		•	•					•	•		
DEN - 1	AUC UUG	ACC	ACC UUG UGG	UGG	GAG GGC	CGC		SSS			חחח	ngg		ACG 1	ACC 1	AUA (300		JCC
DEN-2		A	:	•	Α	Α		Α	ָט						:		Α		A : .
DEN-3	A ACA	Α	ດ.ດ	:	A :	A :	UCA	D.:	ຶ່	Ä		:	:		r U		n:	D	:
DEN - 4	7480	•	:	:	:	:		:	•		:						•		:
DEN-1	ACC GCC	AAC	AAC AUU UUC 1	UUC	AGG	3GA	AGU	UAC	OUG (929	GGA (GCU (GGA	CUG	GCU 1	000	JCA (CAC	AUA
DEN-2	. UG U	:	:	D	Α	ტ		:	•	ن :							ນ:		ტ
DEN-3	o. ug	:	ບ:	D	A :	r :		D ::	ď.	A.							D		o.
DEN - 4	7540	:	:	•	:	:		:	•	:							•	:	
DEN-1	AAG AAU	GCA		CAA ACC	CCU	AGG	AGG												
DEN-2	טיי אייי	C A			A . A	A : c	A :												
DEN-4	4		; ;	:	• • • •		¢ :												

COMPARISON OF THE NS5 NUCLEOTIDE SEQUENCES OF DENGUE VIRUSES

5 64	ACU GGG ACC ACA GGA GAG ACA CUG GGA GAG AAA UGG AAA ACA CAG UUA AAC		UCA GAA UUC AAC ACC UAC AAA AGG AGU GGG AUU AUG GAG GUG GAC AGA UCC GAA GCC	AGU U.C.G .U G .AA A C.CA A A UU A AB A A	AAGU G.A GAGUAAA C.AA G A.U		GGA UUG AAA AGA GGA GAA ACA ACC AAA CAU	A.C C A.C C C G GA. C.C C C C. U A	G A	.CC C GAUG UCU .AUGGAUG U.C AGUG A.C	UUU GUG GAG AGG AAC CUC GUG AAA CCA GAA GGG AAA GUC AUA GAC CUC GGU UGU GGA	D. D	C A A.G C . UU C A . G U.A C	A U A GGG A.G A G A G U G U U	GGC UGG UCA UAU UAU UGU GCU GGG CUG AAG AAA GUC ACU GAA GUG AAG GGA UAC ACA		 A U AUGG ACAC CG AG	GGA CCU GGA CAU GAG GAA CCU AUC CCA AUG GCG ACC UAU GGA UGG AAC CUA	A C A C U.A A G U G	CACAA G.AU U.UAC C AC	\dots	UCU GGA AAA GAU GUA UUU UUC ACA CCA CCU GAG AAA UGU GAU ACC	AG GUUCUCCCAAG N.A U.G U	AGG C.JAU CUGAG A U	AG GUUC U.GC .AAC A.A C GUGCCGC
NS5 7564	DEN-1 GGA DEN-2	DEN-4		DEN-2A	9 0	169		DEN-2 A			7753 DEN-1 UGG	-2	-3	-4	-	7	DEN-4A 7879	٠.	-2	DEN-3C	4,	-1	2	ران ،	

UCU CCG AAAAAA .G U.U	CUC AGA GGA U.G .AC AAUA .A. AAC TCT TC.	ACU CUG GAG CAA AUG .AA AA AC. C.A CAC U.AA AG. C.A GAG A. C	UCC ACC CA	ACA UCC AGA AUG UUA CUG AAU CGA UUC .UUAGG A.UC A GU U. C C A U. C C A	GUG GAC UUA GGC GCU GGA ACA AGA CAU GUG GCA GUG GAA CCA GAG GUA GCC AAC 1	AUU GGC CAG AGG AUA GAG AAU AUA AAA AAU GAA CAC AAG UCA ACA UGG CAU UAUAAAAG C.AGU G.A A UCGG G.AA A.AGGC G.GGUGUGG AGA C.U C CGA U.G C G.AGCA GAC	AAU CCA C.C U
UCC UCUGAUA	ני	d . th .	AAU UCC	AUG ACAUU GUCA	GAC GUG uu u		GAC AAU C.CA
GGU GAG UGGAA 8068	AAU	3UA A A	AGA C	A.A.C.	AGA .C. .A.	GAU C ACA	GAA C
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4

CCC u	GAG	GCC A G	UUC	AAC A G	CUU C	UUVA C C.U	UUU C .AC
AUC G G.G	AAA 	ACA G	GAG A 	UGG	GAG A.U A	AAA	CGC
		GUG A.C A.C	GAG A A	CAA A GG.	AGG A c.u	AAA G 	GCA
GAU C 	GUG U	GAG A.A ACC	AGAG	AA C	GAG A A	GAG	GGA
ngg	AGA C.C	A UG	ACA C	GAA G	AGA G 	AGA C.U	CUG
CCA C	CAG A	AUU C.G G	UGC U	GAUA	CAC G G	AAG A A	UGG
AAA 	CAA	CAA A A.G	AUC G U.A C.G	GUU AC. ACA CAG	GUG U 	GGG A C	A UG
ACA	GGA 	GCA AAG AG. CG.	AGA . GG	UUC	CUC G	A UG	UAC
CUCG	UUC u u	A C A	CCC	GUG A.A C	GAU G A.A	A UG	UGG
00.00 00.00 00.00	CCC	GGC 	AAA .C. .G.	GCA C U CG.	ngg	AA C	AUA
AGA . A. . A.	ACA u .uc	CGA GA.	AAA G	GGA G C	DDD	UAC 	GCA C U
GUG C A	ACC G A G.A	AAA G .UG	AA C A G	AUA U.G	AGG GAA C	GUC G U	CGU A.A A.G
GUG	GAU C	GCA C.G C.C	AGA . AG . GG	GCA C C	gaa agu agc	 	AGU C
GGC A	ACUA	AGA GA. G	UCC GGA GGA GGA	GCA	GAC U 	ACG A .GC U	GGA U C
AAU C	AUG	CCA . A.	CUU A G	AA CU	GAA	GCC . AA . GA	AAA
GUG A.A	GCU A	ACA C	UUC GAA AC. C.C	UCA AGC A.C	GUG U	ngn	GCA
AUG	AUA G G	CGC A.A A.G	GGU AAA A.G	AGG A A	GCA u u	AAA G	AAG A
ncn 	CAA CAA CAA	ACG 	ngg	GUU G C	GAA G .cu	GGA 	GGA
UCA	ACA	GAC 	UUA C.U C.G	AAG	AAA CGU .G.	CAG G.A .U. G.A	000
GCC	GUC	999 900 900 900	nGG	AGA .AG	GCA	AAA CUU	GA G
UCA	, ,	AAA	1 a	ACA 	UCAG AGU	4 4	0 0
IN-1 IN-2 IN-3 IN-4	EN-1 EN-2 EN-3 EN-4	IN-1 IN-2 IN-3	N - 1 N - 2 N - 3 N - 4	EN - 1 EN - 2 EN - 3 EN - 4	N - 1 N - 2 N - 3 N - 4	IN-1 IN-2 IN-3	IN-1 IN-2 IN-3 IN-4
DEN DEN DEN DEN	DEN DEN DEN	DEN	DEN DEN DEN	DEN DEN DEN DEN	DEN DEN DEN	DEN DEN DEN	DEN DEN DEN

CUA GAG UUC GAA GCC CUU GGU UUC AUG AAU GAA GAU CAC UGG UUC AGU AGA GAG AAU UCA CUC U	GGG GGA AAU AUG UAU GCA GAU GAU ACA GCC GGA UGG GAC ACA GCA CC A AA GCC U C AA GCUA	CAG AAU GAG GCU AAA AUC ACU GAC AUC AUG GAG CCU GAA CAU GCU A.AA .AA .UG G.AA A CAA GGAC AAGAAA C.G CAGC C AGGAC CUGGA CAGCUC C.CC AAG	UUU AAG CUG ACC UAC CAACA U.AGCAA	AUG GAU GUU AUA UCC AGA CGU GAC CAG AGA GGA AGU GGA CAG GUC GGA ACU UAU A C G C C	ACU UUC ACC AAU AUG GAG GUC CAA CUA AUA AGA CAA AUG GAG UCU GAA A U G G G G G A G G B<	GAU GAC AUG CAG AAC CCA AAA GGU UUG AAA GAA .U. C.G CAC .UCA GUC .C. GAA GAA .UC .CU .CA C C.U CCG C.A G.G A.G C C.U CCG C.A G.G A.G
DEN-1 CUA DEN-2 U DEN-3U DEN-4G 907: DEN-1 AGU DEN-2 DEN-3	DEN-1 CCG DEN-2 GAA DEN-3C DEN-4 GAU	DEN-1 CUU DEN-2 U.A DEN-3G DEN-4G	DEN-1 AUU DEN-2A DEN-3A DEN-4	DEN-1 GUG DEN-2A DEN-3A DEN-4A	DEN-1 AAT DEN-2 DEN-3 DEN-4	DEN-1 CAA DEN-2 AGC DEN-3 A.G

GAA GAC AAC CCU AAU AUG ACU GAC AAG ACU CCA GUC CAU UCG UGG GAA GAU AUA CCU UAC CUA CAG UGG GAAA G G.AA	GGG AAA AGA GAG GAU UUG UGG UGU GGA UCC CUG AUU GGA CUU UCU UCC AGA GCC ACC UGG GAC .A AC CAACA UGG A.A AGGU C. .A AC CAAC A UGG A.A AGGU	AC AUU CAC ACG GCC AUA ACC CAG GUC AGG AAC CUG AUC GGA AAA GA CAAA AUAA UCCUACU A .UAAC CAAGA .GCACU	10207 10207 10207 UAC AUG CCA GUA AUG AAA AGA UAC AGU GCU CCU UCA GAG AGU GAA GGA GUU CUG UCC3UA AGG GAA GAGA GAG .CUC TGGGAAGG U UCGGUG AAGG GAG UCAGCC A.T TGG AAG.	CAACAACAACAAAGGCUAUUA.CUCAUGACAAGGC.AAAAGUCAGGUCGGAUCAAGCCAUAGUACGGAAAAAACUAUGCUACCUGUGAGCCCCGUC AGGAGGUGUC.G.C.ACC.UAAGCCACAGUACGGAAGAAGCUGUGCAGCCCUGUGAGGCCCCUGCCAAGGACGUUAAAAG	GAAGUCAGGCCACUUGUGCCACGGUUUGAGCAAACCGU GCUGCCUGUAGCUCC GCCAAUAAU GGGAGGC CAAGGACGUUAAA.GAACAGG.CA.CAC.AAUGC.ACAAUGAGUA.AGUGCAUGGCUCCACCU.AA.G AACAAAACAAAA	GUAAUAAUCCCCAGGGAGGCCAUGCGCCACGGAAGCUGUACGCGUGGCAUAUUGGACUAGCGGUUAGAGGAGCCCCUCCCAUGTAAAAU.UCAAAUACGGC	CACUGACAAAACGCAGCAA AA GGGGGCCCGAAGCCAGGAGCUGUACUCCUGGUGGAAGGACUAGAGGUUAGAGGAGAGAU
DEN-1	DEN-1	DEN-1	DEN-1	DEN - 1	DEN-1	DEN-1	DEN-1
DEN-2	DEN-2	DEN-2	DEN-2	DEN - 2	DEN-2	DEN-2	DEN-2
DEN-3	DEN-3	DEN-3	DEN-3	DEN - 3	DEN-3	DEN-3	DEN-3
DEN-4	DEN-4	DEN-4	DEN-4	DEN - 4	DEN-4	DEN-4	DEN-4

DEN-1 CCCCCCCAACAC AAAAACAGCAUAUUGACGCUGGGAAAGAC DEN-2 A. A. DEN-3 GCGCCAACAC DEN-4 10653 DEN-1 GCGCCGCAAGAUGGAUUGGUGUUGAUCCAACAGGUUCU DEN-2 A. AG. A. DEN-2 A. AG. A.	CCCCCCCAACAC AAAAACAGCAUAUUGACGCUGGGAAAGACCAGAGAUCCUGCUGUCUGCAACAUCCAGGCACAGA
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FIGURET

NS1 DEN-1 DEN-2 DEN-3 DEN-4 DEN-4 DEN-2 DEN-4 DEN-4 DEN-4 DEN-4 DEN-1 DEN-1 DEN-1 DEN-1 DEN-1 DEN-3 DEN-3 DEN-4 DEN-1 DEN-3 DEN-1		THE DEDUCED NS1 AMINO ACID SEQUENCES OF DENGUE VIRUSES VINWKG KELKCGSGIF VTNEVHTWTE QYKFQADSPK RLSAAIGKAW EEG VS. N. VDN. PB. S K.AS. Q. H VS. S. VDN. PB. A AS. LN. H KD. NVAN. RGGHDLTVVA GDVKGVLTKG KRALTPPVSD LKY LLL. A. I. NDIK. V CIT. EQ. T. QPME TPEARN STFLIDGPDT SECPNERRAW NSLEVEDYGF GMFTTNIWMK FRE ST.SH. Q. S. L. S. P. SAS. VW. I. L. L. A.TQ. S. I. S. P. SAS. VW. T. LL. L. SAAIKD QKAVHADMGY WIESSKNQTW QIEKASLIEV KTCLWPKTHT LWS NR. NA. D. KM. F. S. H. S. H. S. H. S. H. S. H. S. S. S. S. S. H. S. S. S. S. S. S. H. S.	WCED NS1 AMINO ACID SE KELKCGSGIF VTNEVHTWTE TTNELNYVLW EGGHDLTVVA P. HI S NEVK.IMT A. I. NDIKV STFLIDGPDT SECPNERRAW Q. E. A. N. S.I. S. P. SAS OKAVHADMGY WIESSKNQTW NR. AL.D. ER ESQHNYRQGY ATQTVGPWHL V. P. H. A.	QYKFQADSPK QYKFQADSPK CI.PE.S CI.IMQA. CIT.EQ. NSLEVEDYGF VW. VW. QIEKASLIEV KL. KL. KL. CI.NY CI.NY	QUENCES OF DENGUE VIRUSES QYKFQADSPK RLSAAIGKAW EEGVCGIRST PE.S K.AS.Q.HI.V PE.A .AS.LN.H KD GDVKGVLTKG KRALTPPVSD LKYSWKTWGK .CI.IMQA .S.R.QPTE CIT.EQ. T.QPME CIT. PQT.QPME CIT. PQT.QPME VWV.L.L. L.VYTQL. VWVL. L.VYTQL. QIEKASLIEV KTCLWPKTHT LWSNGVLESQ KM. F. S.H.S GKLEIDFGEC PGTTVTIQED CDHRGPSLRT GKLEIDFGEC PGTTVTIQED CDHRGPSLRT GKLEIDFGEC PGTTVTIQED CDHRGPSLRT M.DF. EVVTGN.	EEGVCGIRST N
DEN-1 DEN-2 DEN-3 DEN-4	TTASGKLVTQ	TTASGKLVTQ WCCRSCTMPPI.EL TVIHEL		•		T. S

COMAPRISON OF THE DEDUCED NS2a AND NS2B AMINO ACID SEQUENCES OF DENGUE VIRUSES

CAIILGGLTW MDLLRALIML VTL.T.NMSF R.G.VMV.V VLLLS.QIRGMAHTI	TALMVIGMAM TTVLSIPHDL LMMATI.L LSQSTETI NL.LGV.L AAT.RL.E.I	LVMAWRTIMA VLFVVTLIPL .QNKVSCT I.AA.SVSTVATL I.AGIS.L.V	WLLNEGIMAV GLVSLLGSAL PAMI.A.S. PVI.A.S. PV	DEMADITGSS PIIEVKQDED EDQ.EISLSITIS E.E.EQV. HNLMITV.D.	YMWQVKTQR .L.EK HTKQ
	VFLRKLTSRE TLLLK. LLFK.	SLTFIRSTMP LAILCVPNAVI CONTIFT	GAQALPVYLM TLMKGASRRS WLLNEGIMAV .LNPTAIF.TRTSKKPAVPPLFIF S.KDTLKPV	SLEKAANVQW E.R.D.K. TVD.T.	TSMITLLVKL DLITVSGLYP LAIPVTMTLW YMWQVKTQR EQTL.I.IRT G.LVIVF. VSI.AAAL.EK ENIL.V.L.T A.LIIF. YSALV. HTKQ
ECLRRRVTRK HMILVVVITL.MTGT.AL.AVSF.VM.GKFGKAG.LF.F	FKMSPGYVLG VR.TFAA. IQ.FLA	DNTQVGTLAL EKY.LAVTIM ETY.LW.ALV	GAQALPVYLM .LNPTAIF.T .VPPLFIF	LKNDLPLASP MVAGGLLLAA YVMSGSSADLI.MTG. LTVCLT.RV.M.G. LI.CIT.T	DLITVSGLYP G.LVIVF. A.LIIF.
ETFSM GLLCLTLFVE DNLGMAL. DN.TVAILF.	GRIGG QIH LAIMAV DDM GVTYLL.A D.M.M GLTYLI.T	LILLKIVTQF MMVRNM .MALI	HWVEITALIL . IPLALTIK . LPM.VAAM	MVAGGLLLAA LTVC LI.C	TSMITLLVKL DLITVSGLYP EQTL.I.IRT G.LVIVF. ENIL.V.L.T A.LIIF.
NS2a GQGTSETFSM .H.QIDNL KVDN.T.	GDTMSGRIGG C.A.TDDM G.SNA.D.M.M G	MELIDGISLG L.T.ALA EQMANA	CRTSCLQKQS LLSQAD .QS.SMR.TD	LKNDLPLASP I.MTG. V.M.G.	GSFSIRDVEEMMNETMR.K.D.T
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

COMPARISON OF THE DEDUCED NS3 AMINO ACID SEQUENCE OF DENGUE VIRUSES

DQYVFSGDPLIYM.EI.M.Q	RKTFVELMRR	KL PRWLDAR K	
RIGRNPAQEDKN.N .VQK.N	DGEFRLRGEQ	EIWTREGEKKKKKK	
TPASAAQRRG . HS . V	GPEREKTQAI EVD ESA	NQILEENMEV	
RVILAGPIPV	TPEGIIPTLF	REWCFTGERN .RD.IKKD	
KPVILTCGPE	EAKMLLDNIY	VASAGISYKDAEN.AEK.T.	FKEFASGRK
GRVIDPRRCL EM D	KNDEDHAHWT ECK NK	GDLPVWLSYK VASAGISYKD REWCFTGERN NQILEENMEV EIWTREGEKK KL PRWLDARA.RAEN.ARD.IKVKRKAHE.K.TKDDKRR	VYSCPLALKD I.DE T.DE
		DEN-1 DEN-2 DEN-3 DEN-4	

COMPARISON OF THE DEDUCED NS4a AND NS4B AMINO ACID SEQUENCES OF DENGUE VIRUSES

		.				
SLETLMLVAL TL.LT. TML.LG.	LEFFRMVLLILI M	ETTILDV QESESNI VVSP.SY	QAAVLMGLGK GWPLHRMDLGTSKIHIVDISK	IMKNPTVDFI	ILTLWEGNPG S TS	
YLHALNELPE S .RVE	QPQWIAASIIH PLSA.V	TDFGFYQVKT K.L.LGSIT. R.L.MSKEPG	QAAVLMGLGKTD.	REAQKRTAAG	CEVLTLATGPA	
LHTTERGGRA	ASGLLWVAEIIY.Q. SMM.DV	ILTIIGLIAA NEMGLIEKTKVVAATMN.FLLAAIVL.TTH	ANLSLAAIAN V.VT V.V	IGPGLQAKAT	LLLMRTTWAF V.ML SL	KNAQTPRR TTNT .SVG.GK.
AKLALDNIVM .RDALV TRNL	LSMGLITIAV MTL.MCC.IT T.ICVIA	ILTIIGLIAA VVAATM. LAAIV	MLRHTIENTSSS.	LGMLLVHYAI .LL.VA VLL.VT	VMLLVLCAGQ	AGAGLAFSLIL.IMLLIM
ASLPTYLSSR GRFMTQK GRV.SH.AH.	FFMQGKGIGK .L.S .LIS	DNQLIYVILTTIAA.VIG	YAVATTILTP	QVNPTTLTASIALI.A	DPKFEKQLGQ 	YANIFRGSYL H
NS4a SITEDILTEI .LNLIM ALV		PEPEKQRTPQ	DLRPASAWTL	VPLLAMGCYSI	TVIDLEPISYP. MTD.VI.	R FWNTTIAVS
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

COMPARISON OF THE DEDUCED NS5 AMINO ACID SEQUENCE OF DENGUE VIRUSES

ET TKHAVSRGTA DSI .HS. SK MSS	GH EEPIPMATYGHSVS	EP WLRGN QFCI .N .NN.TKN	VS AVNWTSRMLL S.HI.H.I S.V.L	IKS TWHYDEDNPY ET SQ SDE EQ	FG QQRVFKEKVD	AIG AVFVDENQWN L. I.INK.K MTED
AKEGLKRGETISAD.SK	KGYTKGGPGHL R	TLRVLKMVEP INLN	VSCGTGNIVSNAS I.N	IENIKNEHKSQET .KRES. LQRLQEE	IAMTDTTPFG N.H ML.VAIH	F RKVRSNAAIG K.T.M.
SGIMEVDRSE QTL TT.	CAGLKKVTEV .GN.R M.TL.N	SPNPTIEEGR	SRNSTHEMYW	VANLDIIGQR IPT.K. TP.M.V.E.	PWDVI PMVTQ	PRICTREEFM
SKSEFNTYKR GEQIK .RKDLK DRKEE	GCGRGGWSYY	CTLLCDIGES .I	HGGMLVRNPL Y.A	GRRHVAVEPE .T.NIGI.S. .TNA	VNGVVRLLTKK	LWGFLSRNKKKE.GKK.TRT.GR
EKWKTQLNQL SRA. KK	VKPEGKVIDL .IVC. .IR	DVFFTPPEKC CY L.YK.T.QV	VETLEQMQRK I.KM.TL I.H.RL	TYERDVDLGATS	VKPSGSASSM T.QT .AT	AQIMEVTAKW KKL.LIE. RKVIE. RMV.TTN.
NS5 GTGTTGETLGNI SQ	KLRWFVERNLMM	WNLVKLHSGKR.QVYV	KILNPYMPSV .VST	NRFTMAHRKPK.KA	KTWAYHGSYE	TRTPRAKRGTQEP.EPMP
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4

SRAIWYMWLG E	YADDTAGWDTI	DVISRRDQRG .I .IKE	CGVDRLKRMA V.KE.S K.E	SHHFHQLIMKI.E.VE AL.KTF	RDLRLAANAI	VHSWEEVPYL .E .TTD	YSAPSESEGVL FRREE.EAW FRKEEAIW
KLGEFGKAKG	DISKIPGGNM .VE.A. E.D.KD.DL.	QRPAKNGTVM : TPR TPK L TPR L TPR . A	KERVEKWSKE IAVON .LAR EKKITQ.LET	WNDWQQVPFC T .H	QMWQLMYFNR TH. SH.	ENPWMEDKTP D	YVDYMPVMKR .TS FLR
VYNMMGKREK	GLHKLGYILR	LTYQNKVVTL	QDDMQNPKGL SIQHLTVTEE KA.LEHP.	DIPOWEPSKG	ETACLGKSYA	LSVWNRVWIE AKQ	QVRNLIGKEESH
LHKQGKCATCLECLGS	ENSLSGVEGE GV.	HALLATSIFK .KKEARQNA	RQMESEGVIT .GGIFK GLS	FLNDMGKVRK AA	VSQGAGWSLR IK I	HHQWMTTEDM K.E.N	WDENIHTAIT AK.O.N AQ.L.O
FWDLVHREREEDNKD	FMNEDHWFSR \cdot	AKITDIMEPE EHV.NHG. EQQ.D DLEQ.A.H	TFTNMEVQLI	LDERFGTSLLDASA.T I.DANA	NQDELVARARIG PIG	TSRTTWSIHA I	LIGL SSRAT
SAKEAVEDERRSRAE	ARGLEFEALG FP	RITEDDLQNELK	SGQVGTYGLN	ISADDCVVKP G	DGREIVVPCR .C.VL KLQ	CSAVPVDWVP	GKREDLWCGS
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

DEN-1 GAA GAC AAC CCU AAU AUG ACU GAC AAG ACU CCA GUC CAU UCG UGG GAA GAU AUA CCU UAC CUA DEN-2 C A G UGG GAA A G G.A A	DEN-1 GGG AAA AGA GAG GAU UUG UGG UGU GGA UCC CUG AUU GGA CUU UCU UCC AGA GCC ACC UGG GAC DEN-2AAC CAACCA UGG A.A AGGUCC DEN-3AGAC CAACACAUC A	DEN-1 GAG AAC AUU CAC ACG GCC AUA ACC CAG GUC AGG AAC CUG AUC GGA AAA GAG GAA UAC GUG GAU DEN-2 ACAAAAAAA	UG CCA GUA AUG AAA AGA UAC AGU GCU CCU UCA GAG AGU GAA GGA GU UCCGUA AGG GAA GAGA GAG .CU	DEN-1 CAACAACAACCAAAGGCUAUU DEN-2A.CUCAUGACAAGGC.AAAGUCAGGIUCGGAUCAAGCCAUAGUACGGAAAAACUAUGCUACCUGUGAGCCCCGUC DEN-3 AGGAGGUGUC.G.C.ACC.UAAGCCACAGUACGGAAGAAGCUGUGCAGCCCUGUGAGGCCCCUGCCAAGGACGUUAAAAG DEN-4	DEN-1 GAAGUCAGGCCACUUGUGCCACGGUUUGAGCAAACCGU GCUGCCUGUAGCUCC GCCAAUAAU DEN-2 CAAGGACGUUAAA.GAACAGG.CA.CAC.AAUGC.ACAAUGAGUA.A GUGCAUGGCUCCACCU.AA.G DEN-3 AACAAAA DEN-4	DEN-1 GUAAUAAUCCCCAGGGAGGCCAUGCGCCACGGAAGCUGUACGCGUGGCAUAUUGGACUAGCGGUUAGAGGAGCCCCUCCCAU DEN-2 GTAAAAU.UCAAAUAGGCU. DEN-3 C.GUAC.G.UUGCAAAGUACUGGCAU.	DEN-1CACUGACAAAACGCAGCAAAA GGGGGCCCGAAGCCAGGAGGAAGCUGUACUCCUGGUGGAAGGACUAGAGGUUAGAGGAGADEN-2U
					ממממ	ממממ	000

CCCCCCAACAC AAAAACAGCAUAUUGACGCUGGGAAAGACCAGAGAUCCUGCUGUCUGCAACAUCCAGGCACAGA A. A. A. A. C. C. AU 10653 3'end length (387) A. AG. A.	CAUAUUGACGCUGGGAAAGACCAGAGAUCCUGCUGUCUCUGCAACAUCAAUCCAGGCACAGA	3'end length (387) (455) (448) (385)
CCCCAACAC AAAACAGCAUAUUGA(A.A	CGCUGGGAAAGACCAGAGAUC	10653 3'end
71 <i>[1</i>]	CCCCCCCAACAC AAAAACAGCAUAUUGA	acccccaagauggauugguguuguugau aag.aaaaa.

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COMPARISON OF THE DEDUCED NS1 AMINO ACID SEQUENCES OF DENGUE VIRUSES

NS1

IRST V 	TWGK	EVCD VF QL	LESQ E D	SLRT	
EEGVCGI	LKYSWKTWGK	FREGSSEVCD LKQDVF LVYTQL	LWSNGVLESQ	CDHRGPSLRT.GN	TA
DSGCVINWKG KELKCGSGIF VTNEVHTWTE QYKFQADSPK RLSAAIGKAW EEGVCGIRSTVSN	KRALTPPVSDS.R.QPTETQPME	GMFTTNIWMK .VI.LVL.	KTCLWPKTHT .S.HS TS	PGTTVTIQED EVVT EV.S.N	KEENMVKSQVL.N.L.
QYKFQADSPKPESPES	GDVKGVLTKG .CIIMQA. .CITEQ.	SECPNERRAW NSLEVEDYGF GMFTTNIWMK ANVI.L. PSASVWL.	HRLMSAAIKD QKAVHADMGY WIESSKNQTW QIEKASLIEV SKNRAL.D KMFV. ERQGS. KL	MLIPKEYAGP FSQHNYRQGY ATQTVGPWHL GKLEIDFGEC PGTTVTIQED .INFVPHAMDF. EVVT SLIPHAL.NY. EV.S.N	TTASGKLVTQ WCCRSCTMPP LRFLGEDGCW YGMEIRPLSE KEENMVKSQVI.ELYRKL.N.LTVIHELYMINLA
VTNEVHTWTE I.DN	EGGHDLTVVA (.NEVKIMT .NDIKV		WIESSKNOTWAL.DQGS.	ATQTVGPWHL HA HA	LRFLGEDGCWYR
KELKCGSGIF	ITNELNYVLWPHI.S .AI	STFLIDGPDT 2E. S.IS.	QKAVHADMGY NR ER	FSQHNYRQGY VP IP	WCCRSCTMPPL
DSGCVINWKGVSN .MVS.S.	TRLENVMWKQ	AKIFTPEARN (MLST.SH. (V.A.TQ.	HRLMSAAIKD SKV	MLIPKEYAGP INF SL	TTASGKLVTQ WI.ETVIHE .
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

COMAPRISON OF THE DEDUCED NS2a AND NS2B AMINO ACID SEQUENCES OF DENGUE VIRUSES

MDLLRALIML R.G.VMV.V RGMAHTI	TTVLSIPHDL LSQSTETI AAT.RL.E.I	VLFVVTLIPL I.AA.SVS I.AGIS.L.V	GLVSLLGSAL	PIIEVKQDEDLSITIS HNLMITV.D.	
ECLRRRVTRK HMILVVVITL CAIILGGLTW MDLLRALIML.M.TGTAL.AVSF VTL.T.NMSF RG.VMV.V.VM.GKFGKAG.LF.F VLLLS.QI RGMAHTI	TALMVIGMAM TTVLSIPHDLLMMATI.L LSQSTETI	LVMAWRTIMA .QNKVSCT .TVATL	WLLNEGIMAV GLVSLLGSAL. P. A M. I.A.S. P V I.A.S. P	DEMADITGSS EDQ.EIS E.E.EQV.	YMWQVKTQR .L.EK HTKQ
HMILVVVITL .AL.AVSFAG.LF.F	VFLRKLTSRE LLK. F	SLTFIRSTMP AILCVPNAVI CSNTIFT	TLMKGASRRSRTSKKS.KDTLK	SLEKAANVQW ERD.K. TVD.T.	LAIPVTMTLW VSI.AAA. YSALV.
ECLRRRVTRK .MTGT. .VM.GKFGK.	FKMSPGYVLG VR.TFAA. IQ.FLA	DNTQVGTLAL EKY.LAVTIM ETY.LW.ALV	GAQALPVYLM TLMKGASRRS.LNPTAIF.TRTSKKVPP.LFIF S.KDTLK	YVMSGSSADLLT.RIT.T	DLITVSGLYP G.LVIVF. A.LIIF.
TFSM GLLCLTLFVE NLGMAL. N.TVAILF.	QIH LAIMAV GVTYLL.A GLTYLI.T	LILLKIVTQF MMVRNM .MALI	HWVEITALIL .IPLALTIK .LPM.VAAM	MVAGGLLLAA LTVC LIC	TSMITLLVKL DLITVSGLYP EQTL.I.IRT G.LVIVF. ENIL.V.L.T A.LIIF.
NS2a GQGTSETFSM G: .H.QIDNL KVDN.T.	GDTMSGRIGG QIH LAIMAV F .A.TDDM GVTYLL.ASNA.D.M.M GLTYLI.T.	MELIDGISLG 1 L.T.ALA 1 EQMANA	CRTSCLQKQS LLSQAD .QS.SMR.TD	LKNDLPLASP MVAGGLLLAA YVI.MTG. LTVCV.M.G. LI.CCVGI.C	GSFSIRDVEE MKNE TMR.K.D.T
DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4

COMPARISON OF THE DEDUCED NS3 AMINO ACID SEQUENCE OF DENGUE VIRUSES

FRKTQVGVGI HMEGVFHTMW LGYS.I.A.V YKE.TGV QKE	DKWD KEEDVQVLAI EPRKNPKHVQ TKPGLFKTLTGE.K EG.EL .GRAN. AQ.Q .G.EI.V .FNFM.I.Q.T.	IGLY GNGVVTKSGD YVSAITQAER IGEPDYEVDE VR.AA.T.K SI.DNP.IED VN.GG.A.TNA EPDGPTPEL.	SIVR EALKREIRNL ILAPTR AA EMEEALRGLP AIG.TVVPMK AITVVPMK	TTRL LSSTRVSNYN LIVMDEAHFT DPSSVAARGYMPVPIA.IMPVPI S.A.I	PPQSN SPIEDIEREI PERSWDTGFD WITDYQGKTV A.M.E NS.HE .VFK A.Q.E.DNS.NEFV	SRKT FDTEYPKTKL TDWDFVVTTD ISEMGANFRAS.V.RA NKQNI.
NG3 SGSLWDVPSP AATQKAALSE GVYRIMQRGL A.V PPVG.E.ED A.KIV PETE.EK.Q.IA	ETGRLEPSWA DVRNDMISYG GGWRLGDKWD KGK.IKK.LK.EGE.K NGKN S.KK.LSAQ.Q	GGIGAVTLDF KPSTSGSPII NRKGKVIGLY .TS S.GV DVEIAGEV	DI FRKKRLT IMDLHPGAGK TKRILPSIVR	IRYQTPAVKS EHTGREIVDL MCHATFTTRLIRAM T.TM	ISTRVEMGEA AAIFMTATPP GATDPFPQSN GGTA.A	WFVPSIKAGN DIANCLRKSG KKVIQLSRKT
DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4

GRVIDPRRCL KPVILTCGPE RVILAGPIPV TPASAAQRRG RIGRNPAQED DQYVFSGDPL E M D.E M HS KN.N IYM.E. D M D.E M VV QK.N I.M.Q. KNDEDHAHWT EAKMLLDNIY TPEGIIPTLF GPEREKTQAI DGEFRLRGEQ RKTFVELMRR E C. K N SM. E VD Y A D. NK N A. E SA Y K.DS GDLPVWLSYK VASAGISYKD REWCFTGERN NQILEENMEV EIWTREGEKK KL PRWLDAR A. R B. K. T K.D D K. R. VYSCPLALKD FKEFASGRK I. D E A A T. D E D A A T. D E D A A T. D E D A	GPIPV TPASAAQRRG RIGRNPAQED DQYVFSGDPLMHS	IPTLF GPEREKTQAI DGEFRLRGEQ RKTFVELMRRSM. EVDXADA. ESAYK.DS	TGERN NQILEENMEV EIWTREGEKK KL PRWLDAR D.IKVKRK D	
	L KPVILTCGPE RVIL MD.E	T EAKMLLDNIY TPEG KNN	K VASAGISYKD REWCRAEN.AR	D FKEFASGRK EA EDA
DEN-1 DEN-3 DEN-3 DEN-4 DEN-1 DEN-1 DEN-2 DEN-4 DEN-2 DEN-2 DEN-2 DEN-2 DEN-3 DEN-3				

COMPARISON OF THE DEDUCED NS4a AND NS4B AMINO ACID SEQUENCES OF DENGUE VIRUSES

	77.07						
DEN-1 DEN-2 DEN-3 DEN-4	SITLDILTEI .LNLIMALV	ASLPTYLSSR GRFMTQK GRV.SH.AH.	AKLALDNIVM .RDALV TRNL	LHTTERGGRA		SLETLMLVAL TL.LT. TML.LG.	
DEN-1 DEN-2 DEN-3 DEN-4	LGAMTAGIFL .ATV.G MILL.G.AM.	FFMQGKGIGK .L.S .LIS	LSMGLITIAV MTL.MCC.IT T.ICVIA	ASGLLWVAEIIY.Q. SM.M.DV	QPQWIAASII H PLSA.V	LEFFRMVLLI LI M	
DEN-1 DEN-2 DEN-3 DEN-4	PEPEKQRTPQ	DNQLIYVILTTIAA.VIG	ILTIIGLIAA VVAATM. LAAIV	NEMGLIEKTK N.FL	TDFGFYQVKT K.L.LGSIT. R.L.MSKEPG	ETTILDV QESESNI VVSP.SY	
DEN-1 DEN-2 DEN-3 DEN-4	DLRPASAWTL	YAVATTILTP MLRHTIENTSFVSSSVST	MLRHTIENTSSS.	ANLSLAAIAN V.VT	ANLSLAAIAN QAAVLMGLGK V.V.TTD.	GWPLHRMDLG SKIHI. ISK	
DEN-1 DEN-2 DEN-3 DEN-4	VPLLAMGCYS	QVNPTTLTASIAL.I.A	LGMLLVHYAI .LL.VA VLL.VT	IGPGLQAKAT	NS4B REAQKRTAAG A	IMKNPTVDFIG.	
DEN - 1 DEN - 2 DEN - 3 DEN - 4	TVIDLEPISYP. MTD.VI.		DPKFEKQLGQ VMLLVLCAGQ	LLLMRTTWAF V.ML	CEVLTLATGPA	ILTLWEGNPG .S .TS	
DEN-1 DEN-2 DEN-3 DEN-4	RFWNTTIAUS K	YANIFRGSYL AGAGLAFSLI HL.IM ML.IM TL.IM	AGAGLAFSLIL.IMLIN	KNAQTPRR TTNT .SVG.GK.			

COMPARISON OF THE DEDUCED NS5 AMINO ACID SEQUENCE OF DENGUE VIRUSES

SGIMEVDRSE AKEGLKRGET TKHAVSRGTAQTLIBSTTSAD.SK MSS	CAGLKKVTEV KGYTKGGPGH EEPIPMATYG .GN.RLHS RN.S	SPNPTIEEGR TLRVLKMVEP WLRGN QFCIA. INLNNN.TSV.S	SRNSTHEMYW VSCGTGNIVS AVNMTSRMLLNASSHIH.II.NSVL	VANLDIIGQR IENIKNEHKS TWHYDEDNPY IPT.KQ.ET SQ TP.M.V.EKR.E.SDE KPDMIISGR. LQRLQEEQ	PWDVIPMVTQ IAMTDTTPFG QQRVFKEKVDVN. M	PRICTREEFT RKVRSNAAIG AVFVDENQWNMLI.I.NK.KLKTMTEDLI.SR.QE.QG.T
SKSEFNTYKR GEQIK .RKDLK DRKEE	VKPEGKVIDL GCGRGGWSYY CAGLK .IVC	DVFFTPPEKC CTLLCDIGES SPNPT C	VETLEQMQRK HGGMLVRNPL SRNST I.KM.TL YA	TYERDVDLGA GRRHVAVEPE VANLETS .T.NIGI.S. IP I.KTNA TP.M.	VKPSGSASSM VNGVVRLLTK PWDVI T.QT	AQIMEVTAKW LWGFLSRNKK PRICTKKL.LIEKE.GKK.TMRKVIERT.GRLRKV.TTNAL.GKK.NL
NS5 GTGTTGETLG EKWKTQLNQLNISRASQKK	KLRWFVERNL VKPEG	WNLVKLHSGK DVFFT	KILNPYMPSV VETLEVST. I.KMVT. I.H	NRFTMAHRKP TYERDK.KATTRR.	KTWAYHGSYE VKPSG	TRTPRAKRGT AQIMEQEP.E KKL.LPMP RKV
DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4

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SRAIWYMWLG	YADDTAGWDT	DVISRRDQRG IK IKE	CGVDRLKRMA V.KE.S K.E	SHHFHQLIMKIE.VE	RDLRLAANAI	VHSWEEVPYL .E .TTD	YSAPSESEGVL FRREE.EAW FRKEEAIW
KLGEFGKAKG	DISKIPGGNM .VEA. E.D.KD.DL.	QRPAKNGTVM TPR TPK LTPR.A	KERVEKWSKE IAVQN .LAR EKKITQ.LET	WNDWQQVPFCTHKE	OMWQLMYFNR TH. SH.	ENPWMEDKTP D	YVDYMPVMKR .TS FLR
VYNMMGKREK	GLHKLGYILR	LTYQNKVVTL	QDDMQNPKGL SIQHLTVTEE KA.LEHP.	DIPQWEPSKGQR.	ETACLGKSYA	LSVWNRVWIE AKQ.T	QVRNLIGKEESH
LHKQGKCATC LEC L.GS.	ENSLSGVEGE GV.	HALLATSIFK .KKEA .RQNA .KIKA	RQMESEGVIT .GGIFK GLS	FLNDMGKVRK A	VSQGAGWSLR IK I	HHQWMTTEDM K.E.N	WDENIHTAIT AK.QN AQLQ
FWDLVHREREEDNKD	FMNEDHWFSR .L	AKITDIMEPE EHV.NHG. EQQ.D DLEQ.A.H	TFTNMEVQLI	LDERFGTSLLDASA.T I.DANA	NQDELVARARIG PIG	TSRTTWSIHA I	LIGL SSRATT T
SAKEAVEDERRSE	ARGLEFEALG FP	RITEDDLQNELK	SGQVGTYGLN	ISADDCVVKP G	DGREIVVPCR .C.VL .KLQ	CSAVPVDWVP	GKREDLWCGS
DEN-1 PTN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN - 1 DEN - 2 DEN - 3 DEN - 4	DEN-1 DEN-2 DEN-3 DEN-4	DEN-1 DEN-2 DEN-3 DEN-4